Fig. 1

SEQ ID NO: 1

GAATTCCGGGCGGCGGCGGCCAAACTCCGCGGCGTCCCCAGGGCGCCA

CGTTCGCCCTCGCCGCCGCGGCCGCGCTGCTCTTCGGCTCCGACATGG

AAGATGGACCTTCTAATAATGCGAGCTGCTTCCGAAGGCTGACCGAGT

GCTTCCTGAGCCCCAAAGCTAATGGTATAGATATAGAAGTCATCCACA

GAGATGTTACAGTTGAAGAGATGGGGGTAGAGAAGACTTTGAAGGAAA

AGAATGTAGAATGAGGATAGAAGAGAGGAAATCCCAACATTTAACAGG

TTTGACAGATGAAAAGTGAAGGCATATCTTTCTCTTCACCCCCAGGT

ATTAGATGAATTTGTATCTGAAAGTGTTAGTGCAGAGACAGTAGAGAA

ATGGCTGAAGAGGAAGAACAACAAATCAGAAGATGAATCAGCTCCTAA

GGAAGTCAGCAGGTACCAAGATACGAATATGCAGGGAGTTGTATATGA

ACTAAACAGCTATATAGAACAACGGTTGGACACAGGAGGAGACAACCA

GCTACTCCTCTATGAACTGAGCAGCATCATTAAAATAGCCACAAAGC

CGATGGATTTGCACTGTATTTCCTTGGAGAGTGCAATAATAGCCTGTG

TATATTCACGCCACCTGGGATAAAGGAAGGAAAACCCCGCCTCATCCC

TGCTGGGCCCATCACTCAGGGCACCACCGTCTCTGCTTATGTGGCCAA

GTCCAGGAAAACACTGCTAGTAGAAGACATCCTTGGAGATGAACGATT

TCCAAGAGGTACTGGACTGGAATCAGGGACTCGTATCCAGTCTGTTCT

TTGCTTACCAATTGTCACTGCAATTGGTGACTTGATTGGTATTCTCGA

GCTGTATCGGCACTGGGGCAAAGAAGCCTTCTGTCTTAGTCACCAGGA

GGTTGCAACAGCAAATCTTGCCTGGGCTTCAGTAGCAATACATCAGGT

GCAGGTATGCAGAGGCCTTGCCAAACAGACAGAATTGAATGACTTCCT

ACTCGACGTATCAAAAACATATTTTGATAACATAGTTGCAATAGATTC

TCTACTTGAACACATAATGATATATGCAAAAAACCTGGTGAATGCCGA

TCGTTGTGCGCTTTTCCAGGTGGACCATAAGAACAAGGAGTTATATTC

AGACCTTTTTGATATTGGAGAGGAAAAGGAAGGAAAACCTGTCTTCAA

GAAGACCAAAGAGATAAGATTTTCAATTGAGAAGGAATTGCTGGCCA

AGTAGCAAGAACAGGGGAAGTCCTGAACATTCCAGATGCCTATGCAGA
CCCACGCTTTAACAGAGAAGTAGACTTGTACACAGGCTACACCACGCG
GAACATCCTGTGCATGCCCATCGTCAGCCGAGGCAGCGTGATAGGTGT
GGTGCAGATGGTCAACAAAATCAGTGGCAGTGCCTTCTCTAAAACAGA
TGAAAACAACTTCAAAATGTTTGCCGTCTTTTGTGCTTTAGCCTTACA
CTGTGCTAATATGTATCATAGAATTCGCCACTCAGAGTGCATTTACCG
GGTAACGATGGAAAAGCTGTCCTACCATAGCATTTGTACTTCAGAAGA
GTGGCAAGGTCTCATGCAATTCACCCTTCCCGTGCGTCTCTGCAAAGA
AATTGAATTATTCCACTTTGACATTGGTCCTTTTGAAAACATGTGGCC
TGGAATTTTTGTCTACATGGTTCATCGGTCCTGTGGGACATCCTGCTT
TGAGCTTGAAAGTTGTGTCGTTTATTATGTCTGTGAAGAAGAACTA
TCGGCGGGTTCCTTATCACAACTGGAAGCATGCGGTCACTGTAGCACA
CTGCATGTATGCCATACTTCAGAACAATCACACGCTTTTCACAGACCT
TGAGCGCAAAGGACTGCTGATTGCGTGTCTGTGTCATGACCTGGACCA
CAGGGGCTTCAGTAACAGCTACCTGCAGAAGTTCGACCACCCTCTGGC
CGCTCTCTACTCCACTTCCACCATGGAGCAGCACCACTTCTCCCAGAC
TGTGTCCATCCTTCAGTTGGAAGGGCACAATATCTTCTCCACTCTGAG
CTCCAGTGAATATGAGCAGGTGCTTGAGATCATCCGCAAAGCCATCAT
TGCCACAGACCTTGCTTTATACTTTGGAAACAGGAAGCAGTTGGAAGA
GATGTACCAGACCGGATCACTAAACCTTAATAATCAATCACATAGAGA
CCGTGTAATTGGTTTGATGATGACTGCCTGTGACCTTTGTTCTGTGAC
AAAACTGTGGCCCGTTACAAAATTGACGGCAAATGATATATATGCAGA
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TCCTATGATGGACAGAGACAAGAAGGATGAAGTCCCCCAAGGCCAGCT
TGGGTTCTACAATGCCGTGGCCATTCCCTGCTATACAACCCTTACCCA
GATCCTCCCTCCCACGGAGCCTCTTCTGAAAGCATGCAGGGATAATCT
CAGTCAGTGGGAGAAGGTGATTCGAGGGGAGGAGACTGCAACCTGGAT
TTCATCCCCATCCGTGGCTCAGAAGGCAGCTGCATCTGAAGATTGAGC
ACTGGTCACCCTGACACGCTGTCCCACCTACAGATCCTCATCTTGCTT

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CTTTGACATTCTTTTCCTTTTTTTGGGGGGGGTGGGGGGAACCTGCAC
CTGGTAACTGGGGTGCAAACCTCTTCAAGAAGGTAACATCAAATAAAT
AAGTCAAGCAGAGGACTTCCTGCCAATCTCTTCTGTGAGGCATCATAG
ACACTGAGCAACCAGGACCACCCCACGTTCAGAAATCAGCTGGCCAA
GTGACTCCATTTGACTTGCAAACCAGCCTTTTCTAATAGGCTAATATT
GCTGAGGCCTTAAAGGAAATGGACAAAAATTATCCAGAAGGGGTACTT
TTCCATTGTATCTTTCTAATAAGGGTTTAAAATGGTACTATTATGGTA
TTGTACTTGGGCTTTAACATCAATGTTGCTTTGATGTTGTTGGATATA
AATAGGAATTTTACACATTACTATTGTGAATGGTGAATGTTCATGTAT
GACCTACTTGTAATTAACTTGAGTTGTAGTCCACAGCCTCAGGACAAA
TGTCGTTGAGGTTACAGAGTAAGAAATGATGGCAAAACGTCAAACTCT
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TTCAACCTAAAGGGCTGAGATCAATAAATTTAACACTAGACGAAGTAG
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TTGTTGTTGTCTTGGCAGTTTTACTGAACCCTTTGCATAAAGAACAAA
ATAAAAGCTCGGCATTGTAATTTTTTTAATGGACAAGTCTTATGGATA
CGAGGGGTACATTTTTCATAATGATTCCTTTATATTTTCACTTTGTGT
CATATGCAGAATTTTAGACTCTCATTCACAATGAAAAGTTTATTTTAA
ACATTGTTTAATTAAAATACCATACAGTTCTCTTTTAAACATCAAACC
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TCACTTATCTCTTCCACGTACTGCACGGTCGTATTTCATGAGCTTTCT
GTCCATAGCACAGAAACAGAGCAGAAAGTAGTACAATCATGTTGGACC
TTCTTTCTGTTCTCTTTACTCTTCTCACAGATCAGATCACTCCATAGA
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GTATTACTATACAATGTAAGTGTTTTAAAAAATACGAAAGTAATACTC
TGCACCCCTTCCTACAAAGATGATAAAGCAGTCACTTCTGGCGCATTT
TAATAATTTAAAGATTTTTAGTGCAATGGCACGGTAACCTCCAAACCT
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TAACTTATTCACTTTTCATGTGGCAGGAAACTGGAATATCGCTTTTAA

TAAAATGGAAAAATATGCTTCTACATATTTACCACCATAGGCGTTTTG

TTCATATGAGCCTGGTTTGTGCAAAATTAAATCAGAGGCTTCTACAAC

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TTATTTTAAAATTAACACTATGTGTTCAGTTTTCTTGTGGGCTTCTGA

AAGTTGCCATCTTCCCTCCGTGGAGCTCCATTTGCTATTTTCATTATA

CACTATGAGGTAAAATGTAATAACAAAGAGAGAGAAGTACCACTGTG

GCTAGATATATACACACATATATATATGGATGGATGTAATATATGT

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GTATACATATGTGTATATGATACATACACATACACACACGAGAG

ACAGAAGGAAAGAGAGGAAGAGAGAAGCAAACATGTAGGAAAAAATAT

AAATCAGCCGGAATTC

Fig. 2

SEQ ID NO: 2

MRIEERKSQHLTGLTDEKVKAYLSLHPQVLDEFVSESVSAETVEKWLK
RKNNKSEDESAPKEVSRYQDTNMQGVVYELNSYIEQRLDTGGDNQLLL
YELSSIIKIATKADGFALYFLGECNNSLCIFTPPGIKEGKPRLIPAGP
ITQGTTVSAYVAKSRKTLLVEDILGDERFPRGTGLESGTRIQSVLCLP
IVTAIGDLIGILELYRHWGKEAFCLSHQEVATANLAWASVAIHQVQVC
RGLAKQTELNDFLLDVSKTYFDNIVAIDSLLEHIMIYAKNLVNADRCA
LFQVDHKNKELYSDLFDIGEEKEGKPVFKKTKEIRFSIEKGIAGQVAR
TGEVLNIPDAYADPRFNREVDLYTGYTTRNILCMPIVSRGSVIGVVQM
VNKISGSAFSKTDENNFKMFAVFCALALHCANMYHRIRHSECIYRVTM
EKLSYHSICTSEEWQGLMQFTLPVRLCKEIELFHFDIGPFENMWPGIF
VYMVHRSCGTSCFELEKLCRFIMSVKKNYRRVPYHNWKHAVTVAHCMY
AILQNNHTLFTDLERKGLLIACLCHDLDHRGFSNSYLQKFDHPLAALY
STSTMEQHHFSQTVSILQLEGHNIFSTLSSSEYEQVLEIIRKAIIATD
LALYFGNRKQLEEMYQTGSLNLNNQSHRDRVIGLMMTACDLCSVTKLW
PVTKLTANDIYAEFWAEGDEMKKLGIQPIPMMDRDKKDEVPQGQLGFY
NAVAIPCYTTLTQILPPTEPLLKACRDNLSQWEKVIRGEETATWISSP
SVAQKAAASED

Fig. 3

SEQ ID NO: 3
5'-TTTTCAAGCTCAAAGCAGGATG-3

Fig. 4

SEQ ID NO: 4
5'-ACATGTGGCCTGGAATTTTTGT-3'

Fig. 5

<u>SEQ ID NO: 5</u>

5'-CCCACAGGACCGATGAACCATG-3'